PCT

RAW SEQUENCE LISTING   DATE: 10/26/2004
PATENT APPLICATION: US/10/511,480   TIME: 18:02:21

Input Set : A:\P28268PC.ST25.txt
Output Set: N:\CRF4\10262004\J511480.raw

```
        3 <110> APPLICANT: Medical Research Council
        5 <120> TITLE OF INVENTION: Treatment Methods
        7 <130> FILE REFERENCE: ARDW/P28268PC
C-->    9 <140> CURRENT APPLICATION NUMBER: US/10/511,480
C-->   10 <141> CURRENT FILING DATE: 2004-10-15
       12 <150> PRIOR APPLICATION NUMBER: GB 0208785.6
       13 <151> PRIOR FILING DATE: 2002-04-17
       15 <160> NUMBER OF SEQ ID NOS: 20
       17 <170> SOFTWARE: PatentIn version 3.1
```

Does Not Comply
Corrected Diskette Needed

(PS.I)

ERRORED SEQUENCES

```
      305 <210> SEQ ID NO: 20
      306 <211> LENGTH: 22
      307 <212> TYPE: DNA
      308 <213> ORGANISM: Artificial
      310 <220> FEATURE:
      311 <223> OTHER INFORMATION: Human ribosomal 18S PCR probe
      313 <400> SEQUENCE: 20
      314 tgctggcacc agacttggcc tc                              22
E-->  317 1
E-->  320 1
```

Deleted

RAW SEQUENCE LISTING ERROR SUMMARY   DATE: 10/26/2004
PATENT APPLICATION: **US/10/511,480**   TIME: 18:02:22

Input Set : **A:\P28268PC.ST25.txt**
Output Set: **N:\CRF4\10262004\J511480.raw**

<u>Invalid <213> Response:</u>
**Use of "Artificial" only as "<213> Organism" response is incomplete,
per 1.823(b) of New Sequence Rules. Valid response is Artificial Sequence.**

Seq#:5,6,7,8,9,10,11,12,13,14,15,16,17,18,19,20

VERIFICATION SUMMARY   DATE: 10/26/2004
PATENT APPLICATION: **US/10/511,480**   TIME: 18:02:22

Input Set : **A:\P28268PC.ST25.txt**
Output Set: **N:\CRF4\10262004\J511480.raw**

L:9 M:270 C: Current Application Number differs, Replaced Current Application Number
L:10 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:216 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:13
L:220 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:13,Line#:218
L:235 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:14
L:239 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:14,Line#:237
L:317 M:254 E: No. of Bases conflict, this line has no nucleotides.
M:254 Repeated in SeqNo=20